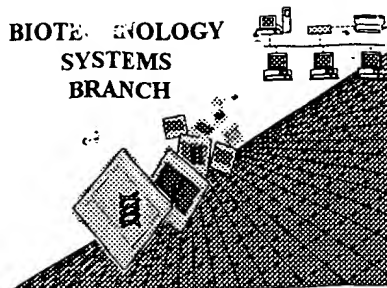


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/613,591B
Source: O/E
Date Processed by STIC: 6/5/2001

RECEIVED

NOV 13 2001

TECH CENTER 1600/2900

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

#19/8W
11-15-01
OIFE

RAW SEQUENCE LISTING

DATE: 06/05/2001

PATENT APPLICATION: US/09/613,591B

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: BOYLE, WILLIAM
 4 LACEY, DAVID
 5 CALZONE, FRANK
 6 CHANG, MING-SHI
 7 SENALDI, GIORGIO
 9 <120> TITLE OF INVENTION: COMBINATION THERAPY FOR CONDITIONS LEADING TO BONE LOSS
 11 <130> FILE REFERENCE: A-378CIP5
 13 <140> CURRENT APPLICATION NUMBER: US 09/613,591B
 14 <141> CURRENT FILING DATE: 2000-07-10
 16 <150> PRIOR APPLICATION NUMBER: US 09/457,647
 17 <151> PRIOR FILING DATE: 1999-12-09
 19 <150> PRIOR APPLICATION NUMBER: US 09/350,670
 20 <151> PRIOR FILING DATE: 1999-07-09
 22 <150> PRIOR APPLICATION NUMBER: US 08/706,945
 23 <151> PRIOR FILING DATE: 1996-09-03
 25 <150> PRIOR APPLICATION NUMBER: US 08/577,788
 26 <151> PRIOR FILING DATE: 1995-12-22
 28 <160> NUMBER OF SEQ ID NOS: 168
 30 <170> SOFTWARE: PatentIn version 3.0
 32 <210> SEQ ID NO: 1
 33 <211> LENGTH: 36
 34 <212> TYPE: DNA
 35 <213> ORGANISM: Artificial
 37 <220> FEATURE:
 38 <223> OTHER INFORMATION: Random cDNA with internal NotI restriction site.
 40 <220> FEATURE:
 41 <221> NAME/KEY: misc_feature *h's are not at these locations*
 42 <222> LOCATION: (16, 17, 18, 19, 20, 21, 22 and)..(23) *these are the correct locations*
 43 <223> OTHER INFORMATION: N = any random nucleic acid
 46 <400> SEQUENCE: 1
 W--> 47 aaaggaagga aaaaagcggc cgctacannn nnnnt 36
 50 <210> SEQ ID NO: 2
 51 <211> LENGTH: 16
 52 <212> TYPE: DNA
 53 <213> ORGANISM: Artificial
 55 <220> FEATURE:
 56 <223> OTHER INFORMATION: ds oligonucleotide adapter
 58 <400> SEQUENCE: 2
 59 tcgaccacg cgtccg 16
 62 <210> SEQ ID NO: 3
 63 <211> LENGTH: 12
 64 <212> TYPE: DNA
 65 <213> ORGANISM: Artificial
 67 <220> FEATURE:
 68 <223> OTHER INFORMATION: ds oligonucleotide adapter
 70 <400> SEQUENCE: 3
 71 gggtgcgcag gc 12

RAW SEQUENCE LISTING

DATE: 06/05/2001

PATENT APPLICATION: US/09/613,591B

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

74 <210> SEQ ID NO: 4
 75 <211> LENGTH: 18
 76 <212> TYPE: DNA
 77 <213> ORGANISM: Artificial
 79 <220> FEATURE:
 80 <223> OTHER INFORMATION: PCR primer
 82 <400> SEQUENCE: 4
 83 tgtaaaacga cggccagt 18
 86 <210> SEQ ID NO: 5
 87 <211> LENGTH: 18
 88 <212> TYPE: DNA
 89 <213> ORGANISM: Artificial
 91 <220> FEATURE:
 92 <223> OTHER INFORMATION: PCR primer
 94 <400> SEQUENCE: 5
 95 caggaaacag ctatgacc 18
 98 <210> SEQ ID NO: 6
 99 <211> LENGTH: 20
 100 <212> TYPE: DNA
 101 <213> ORGANISM: Artificial
 103 <220> FEATURE:
 104 <223> OTHER INFORMATION: T3 primer
 106 <400> SEQUENCE: 6
 107 caattaaccc tcactaaagg 20
 110 <210> SEQ ID NO: 7
 111 <211> LENGTH: 23
 112 <212> TYPE: DNA
 113 <213> ORGANISM: Rattus rattus
 115 <400> SEQUENCE: 7
 116 gcattatgac ccagaaaccg gac 23
 119 <210> SEQ ID NO: 8
 120 <211> LENGTH: 23
 121 <212> TYPE: DNA
 122 <213> ORGANISM: Rattus rattus
 124 <400> SEQUENCE: 8
 125 aggtagcgcc cttcctcaca ttc 23
 128 <210> SEQ ID NO: 9
 129 <211> LENGTH: 30
 130 <212> TYPE: DNA
 131 <213> ORGANISM: Artificial
 133 <220> FEATURE:
 134 <223> OTHER INFORMATION: PCR primer
 136 <400> SEQUENCE: 9
 137 gactagtccc acaatgaaca agtggctgtg 30
 140 <210> SEQ ID NO: 10
 141 <211> LENGTH: 45
 142 <212> TYPE: DNA
 143 <213> ORGANISM: Artificial
 145 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 06/05/2001

PATENT APPLICATION: US/09/613,591B

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

146 <223> OTHER INFORMATION: PCR primer
 148 <400> SEQUENCE: 10
 149 ataagaatgc ggccgctaaa ctatgaaaca gcccagtgac cattc 45
 152 <210> SEQ ID NO: 11
 153 <211> LENGTH: 21
 154 <212> TYPE: DNA
 155 <213> ORGANISM: Artificial
 157 <220> FEATURE:
 158 <223> OTHER INFORMATION: PCR primer
 160 <400> SEQUENCE: 11
 161 gcctctagaa agagctggga c 21
 164 <210> SEQ ID NO: 12
 165 <211> LENGTH: 21
 166 <212> TYPE: DNA
 167 <213> ORGANISM: Artificial
 169 <220> FEATURE:
 170 <223> OTHER INFORMATION: PCR primer
 172 <400> SEQUENCE: 12
 173 cgccgtgttc catttatgag c 21
 176 <210> SEQ ID NO: 13
 177 <211> LENGTH: 24
 178 <212> TYPE: DNA
 179 <213> ORGANISM: Rattus rattus
 181 <400> SEQUENCE: 13
 182 atcaaaggca gggcatactt cctg 24
 185 <210> SEQ ID NO: 14
 186 <211> LENGTH: 24
 187 <212> TYPE: DNA
 188 <213> ORGANISM: Rattus rattus
 190 <400> SEQUENCE: 14
 191 gttgcactcc tgtttcacgg tctg 24
 194 <210> SEQ ID NO: 15
 195 <211> LENGTH: 24
 196 <212> TYPE: DNA
 197 <213> ORGANISM: Rattus rattus
 199 <400> SEQUENCE: 15
 200 caagacacct tgaaggcct gatg 24
 203 <210> SEQ ID NO: 16
 204 <211> LENGTH: 24
 205 <212> TYPE: DNA
 206 <213> ORGANISM: Rattus rattus
 208 <400> SEQUENCE: 16
 209 taactttttac agaagagcat cagc 24
 212 <210> SEQ ID NO: 17
 213 <211> LENGTH: 33
 214 <212> TYPE: DNA
 215 <213> ORGANISM: Rattus rattus
 217 <400> SEQUENCE: 17
 218 agcgcggccg catgaacaag tggctgtgct gcg 33

RAW SEQUENCE LISTING

DATE: 06/05/2001

PATENT APPLICATION: US/09/613,591B

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

221 <210> SEQ ID NO: 18
 222 <211> LENGTH: 31
 223 <212> TYPE: DNA
 224 <213> ORGANISM: Rattus rattus
 226 <400> SEQUENCE: 18
 227 agctctagag aaacagccca gtgaccattc c 31
 230 <210> SEQ ID NO: 19
 231 <211> LENGTH: 24
 232 <212> TYPE: DNA
 233 <213> ORGANISM: Rattus rattus
 235 <400> SEQUENCE: 19
 236 gtgaagctgt gcaagaacct gatg 24
 239 <210> SEQ ID NO: 20
 240 <211> LENGTH: 24
 241 <212> TYPE: DNA
 242 <213> ORGANISM: Rattus rattus
 244 <400> SEQUENCE: 20
 245 atcaaaggca gggcatactt cctg 24
 248 <210> SEQ ID NO: 21
 249 <211> LENGTH: 24
 250 <212> TYPE: DNA
 251 <213> ORGANISM: Homo sapiens
 253 <400> SEQUENCE: 21
 254 cagatcctga agctgctcag ttg 24
 257 <210> SEQ ID NO: 22
 258 <211> LENGTH: 33
 259 <212> TYPE: DNA
 260 <213> ORGANISM: Homo sapiens
 262 <400> SEQUENCE: 22
 263 agcgcgccg cggggaccac aatgaacaag ttg 33
 266 <210> SEQ ID NO: 23
 267 <211> LENGTH: 33
 268 <212> TYPE: DNA
 269 <213> ORGANISM: Homo sapiens
 271 <400> SEQUENCE: 23
 272 agctctagaa ttgtgaggaa acagctcaat ggc 33
 275 <210> SEQ ID NO: 24
 276 <211> LENGTH: 39
 277 <212> TYPE: DNA
 278 <213> ORGANISM: Artificial
 280 <220> FEATURE:
 281 <223> OTHER INFORMATION: PCR primer
 283 <400> SEQUENCE: 24
 284 atagcgccg ctgagcccaa atcttgtgac aaaactcac 39
 287 <210> SEQ ID NO: 25
 288 <211> LENGTH: 45
 289 <212> TYPE: DNA
 290 <213> ORGANISM: Artificial
 292 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 06/05/2001

PATENT APPLICATION: US/09/613,591B

TIME: 12:57:12

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

```

293 <223> OTHER INFORMATION: PCR primer
295 <400> SEQUENCE: 25
296 tctagagtcg acttatcatt tacccggaga caggagagg ctctt      45
299 <210> SEQ ID NO: 26
300 <211> LENGTH: 38
301 <212> TYPE: DNA
302 <213> ORGANISM: Mus musculus
304 <400> SEQUENCE: 26
305 cctctgagct caagcttccg aggaccacaa tgaacaag      38
308 <210> SEQ ID NO: 27
309 <211> LENGTH: 43
310 <212> TYPE: DNA
311 <213> ORGANISM: Mus musculus
313 <400> SEQUENCE: 27
314 cctctgcggc cgctaagcag cttattttca cggattgaac ctg      43
317 <210> SEQ ID NO: 28
318 <211> LENGTH: 38
319 <212> TYPE: DNA
320 <213> ORGANISM: Mus musculus
322 <400> SEQUENCE: 28
323 cctctgagct caagcttccg aggaccacaa tgaacaag      38
326 <210> SEQ ID NO: 29
327 <211> LENGTH: 24
328 <212> TYPE: DNA
329 <213> ORGANISM: Homo sapiens
331 <400> SEQUENCE: 29
332 tccgtaagaa acagcccagt gacc      24
335 <210> SEQ ID NO: 30
336 <211> LENGTH: 31
337 <212> TYPE: DNA
338 <213> ORGANISM: Mus musculus
340 <400> SEQUENCE: 30
341 cctctgcggc cgctgttgca tttcctttct g      31
344 <210> SEQ ID NO: 31
345 <211> LENGTH: 19
346 <212> TYPE: PRT
347 <213> ORGANISM: Mus musculus
349 <400> SEQUENCE: 31
351 Glu Thr Leu Pro Pro Lys Tyr Leu His Tyr Asp Pro Glu Thr Gly His
352 1          5          10          15
354 Gln Leu Leu
357 <210> SEQ ID NO: 32
358 <211> LENGTH: 21
359 <212> TYPE: DNA
360 <213> ORGANISM: Mus musculus
362 <400> SEQUENCE: 32
363 tcccttgccc tgaccactct t      21
366 <210> SEQ ID NO: 33
367 <211> LENGTH: 34

```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:13

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

L:35 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:47 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:53 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:65 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:77 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:89 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:101 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:131 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:143 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:155 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:167 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:278 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:290 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:486 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:46
L:558 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:48
L:570 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:49
L:582 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50
L:651 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:51
L:663 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:52
L:675 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:53
L:691 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:54
L:707 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:55
L:719 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:56
L:731 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:57
L:765 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:58
L:841 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:63
L:856 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:64
L:871 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:65
L:883 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:66
L:895 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:67
L:909 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:68
L:941 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:71
L:955 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:72
L:969 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:73
L:1048 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:81
L:1060 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:82
L:1072 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:83
L:1084 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:84
L:1096 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:85
L:1108 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:86
L:1120 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:87
L:1132 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:88
L:1144 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:89
L:1156 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:90
L:1168 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:91
L:1182 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:92
L:1196 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:93
L:1208 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:94

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/613,591B

DATE: 06/05/2001

TIME: 12:57:13

Input Set : A:\A-378CIP5.ST25.txt

Output Set: C:\CRF3\06052001\I613591B.raw

L:1220 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:95
L:1232 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:96
L:1244 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:97
L:1256 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:98
L:1268 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:99
L:1280 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:100
L:1360 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:108
L:1372 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:109
L:1384 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:110
L:1396 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:111
L:1408 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:112
L:1420 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:113
L:1432 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:114
L:1444 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:115
L:1463 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:116
L:1477 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:117
L:1491 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:118
L:2302 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:127
L:2752 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:137
L:2904 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:140
L:2916 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:141
L:2928 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:142
L:2940 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:143
L:2952 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:144
L:2964 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:145
L:2990 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:147
L:3002 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:148
L:3014 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:149
L:3026 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:150
L:3038 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:151
L:3050 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:152
L:3062 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:153
L:3074 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:154
L:3086 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:155
L:3098 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:156
L:3110 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:157
L:3122 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:158
L:3134 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:159
L:3146 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:160
L:3158 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:161
L:3170 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:162
L:3182 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:163
L:3194 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:164
L:3206 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:165
L:3218 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:166
L:3230 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:167
L:3242 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:168